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Fig. 1A
SEQ. ID NO:1

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PTRLNSASLKQPYITQNYFPVGTVVEYECRPGYRREPSLSPKLTCLQNLK  150
WSTAVEFCKKKSCPNPGEIRNGQIDVPGGILFGATISFSCNTGYKLFGST  200
SSFCLISGSSVQWSDPLPECREIYCPAPPQIDNGIIQGERDHYGYRQSVT  250
YACNKGFTMIGEHSIYCTVNNDEGEWSGPPPECRGKSLTSKVPPTVQKPT  300
TVNVPTTEVSPTSQKTTTKTTTPNAQATRSTPVSRTTKHFHETTPNKGSG  350
TTSGTTRLLSGHTCFTLTGLLGTLVTMGLLT
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Fig. 1B
SEQ. ID NO:2

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 841 tgattggaga gcactctatt tattgtactg tgaataatga tgaaggagag tggagtggcc
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1441 tgagagtgat tcctttccta aaagtgtaag aaagcataga gatttgttcg tatttagaat
1501 gggatcacga ggaaaagaga aggaaagtga tttttttcca caagatctgt aatgttattt
1561 ccacttataa aggaaataaa aaatgaaaaa cattatttgg atatcaaaag caaataaaaa
1621 cccaattcag tctcttctaa gcaaaattgc taaagagaga tgaaccacat tataaagtaa
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1741 acatgctggt gaaccagggg tgttgatggt gataagggag gaatatagaa tgaaagactg
1801 aatcttcctt tgttgcacaa atagagtttg gaaaaagcct gtgaaaggtg tcttctttga
1861 cttaatgtct ttaaaagtat ccagagatac tacaatatta acataagaaa agattatata
1921 ttatttctga atcgagatgt ccatagtcaa atttgtaaat cttattcttt tgtaatattt
1981 atttatattt atttatgaca gtgaacattc tgattttaca tgtaaaacaa gaaaagttga
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2101 gt
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Fig. 2

SEQ. ID NO:3

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IISGDTVIWDNETPICDRIPCGLPPTITNGDFISTNRENFHYGSVVTYRC 200
NPGSGGRKVFELVGEPSIYCTSNDDQVGIWSGPAPQCIIPNKCTPPNVEN 250
GILVSDNRSLFSLNEVVEFRCQPGFVMKGPRRVKCQALNKWEPELPSCSR 300
VCQPPPDVLHAERTQRDKDNFSPGQEVFYSCEPGYDLRGAASMRCTPQGD 350
WSPAAPTCEVKSCDDFMGQLLNGRVLFPVNLQLGAKVDFVCDEGFQLKGS 400
SASYCVLAGMESLWNSSVPVCEQIFCPSPPVIPNGRHTGKPLEVFPFGKA 450
VNYTCDPHPDRGTSFDLIGESTIRCTSDPQGNGVWSSPAPRCGILGHCQA 500
PDHFLFAKLKTQTNASDFPIGTSLKYECRPEYYGRPFSITCLDNLVWSSP 550
KDVCKRKSCKTPPDPVNGMVHVITDIQVGSRINYSCTTGHRLIGHSSAEC 600
ILSGNAAHWSTKPPICQRIPCGLPPTIANGDFISTNRENFHYGSVVTYRC 650
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GILVSDNRSLFSLNEVVEFRCQPGFVMKGPRRVKCQALNKWEPELPSCSR 750
VCQPPPDVLHAERTQRDKDNFSPGQEVFYSCEPGYDLRGAASMRCTPQGD 800
WSPAAPTCEVKSCDDFMGQLLNGRVLFPVNLQLGAKVDFVCDEGFQLKGS 850
SASYCVLAGMESLWNSSVPVCEQIFCPSPPVIPNGRHTGKPLEVFPFGKA 900
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GSPWSQCQADDRWDPPLAKCTSRAHDALIVGTLSGTIFFILLIIFLSWII 2000
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Fig. 3
SEQ. ID NO:4

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Fig. 4A

SEQ. ID NO:5

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SLIGESTIYCGDNSVWSRAAPECKVVKCRFPVVENGKQISGFGKKFYYKA  250
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Fig. 4B

SEQ. ID NO:6

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1381 agattgcctg ctttccctta aataacactt agatttattg gaccagtcag cacagcatgc
1441 ctggttgtat taaagcaggg atatgctgta ttttataaaa ttggcaaaat tagagaaata
1501 tagttcacaa tgaaattata ttttctttgt
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Fig. 5

Fig. 6A
SEQ. ID NO:7
ATA TAC GAA TTC AGA TCT ATG ACC GTC GCG CGG CCG AGC GTG

Fig. 6B
SEQ. ID NO:8
ACA GTG CTC GAG CAT TCA GGT GGT GGG CCA CTC CA

Fig 7A
SEQ. ID NO:9
ATA TAC CTC GAG TCC TAA CAA ATG CAC GCC TCC AAA TGT GG-3

Fig 7B
SEQ. ID NO:10
ACA GTG ATG CAT TGG TTT GGG TTT TCA ACT TGG C

Fig 7C
SEQ. ID NO:11
ATA TAC ATG CAT CTG ACT TTC CCA TTG GGA CAT CTT TAA AG

Fig 7D
SEQ. ID NO:12
ACA GTG AGA TCT TTA GTG ATG GTG ATG GTG ATG AAT TCC ACA GCG AGG GGC AGG GCT

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Fig. 8A
SEQ ID NO:13

M T V A R P S V P A A L P
L L G E L P R L L L L V L L C L P A V W G
D C G L P P D V P N A Q P A L E G R T S F
P E D T V I T Y K C E E S F V K I P G E K
D S V I C L K G S Q W S D I E E F C N R S
C E V P T R L N S A S L K Q P Y I T Q N Y
F P V G T V V E Y E C R P G Y R R E P S L
S P K L T C L Q N L K W S T A V E F C K K
K S C P N P G E I R N G Q I D V P G G I L
F G A T I S F S C N T G Y K L F G S T S S
F C L I S G S S V Q W S D P L P E C R E I
Y C P A P P Q I D N G I I Q G E R D H Y G
Y R Q S V T Y A C N K G F T M I G E H S I
Y C T V N N D E G E W S G P P P E C
S S P N K C T P P N V E N G I L V S D N
R S L F S L N E V V E F R C Q P G F V M
K G P R R V K C Q A L N K W E P E L P S
C S R V C Q P P P D V L H A E R T Q R D
K D N F S P G Q E V F Y S C E P G Y D L
R G A A S M R C T P Q G D W S P A A P T
C E V K S C D D F M G Q L L N G R V L F
P V N L Q L G A K V D F V C D E G F Q L
K G S S A S Y C V L A G M E S L W N S S
V P V C E Q I F C P S P P V I P N G R H
T G K P L E V F P F G K A V N Y T C D P
H P D R G T S F D L I G E S T I R C T S
D P Q G N G V W S S P A P R C G I L G H
C Q A P D H F L F A K L K T Q T N A S D
F P I G T S L K Y E C R P E Y Y G R P F
S I T C L D N L V W S S P K D V C K R K
S C K T P P D P V N G M V H V I T D I Q
V G S R I N Y S C T T G H R L I G H S S
A E C I L S G N A A H W S T K P P I C Q
R I P C G L P P T I A N G D F I S T N R
E N F H Y G S V V T Y R C N P G S G G R
K V F E L V G E P S I Y C T S N D D Q V
G I W S G P A P Q C I I P N K C T P P N
V E N G I L V S D N R S L F S L N E V V
E F R C Q P G F V M K G P R R V K C Q A
L N K W E P E L P S C S R V C Q P P P D
V L H A E R T Q R D K D N F S P G Q E V
F Y S C E P G Y D L R G A A S M R C T P
Q G D W S P A A P T C E V K S C D D F M
G Q L L N G R V L F P V N L Q L G A K V
D F V C D E G F Q L K G S S A S Y C V L
A G M E S L W N S S V P V C E Q I F C P
S P P V I P N G R H T G K P L E V F P F
G K A V N Y T C D P H P D R G T S F D L

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I G E S T I R C T S D P Q G N G V W S S
P A P R C G I H H H H H H

Fig. 8B
SEQ. ID NO: 14

ATGACCGTCGCGCGGCCGAGCGTGCCCGCGGCGCTGCCC
CTCCTCGGGGAGCTGCCCCGGCTGCTGCTGCTGGTGCTGTTGTGCCTGCCGGCCGTGTGGGGT
GACTGTGGCCTTCCCCCAGATGTACCTAATGCCCAGCCAGCTTTGGAAGGCCGTACAAGTTTT
CCCGAGGATACTGTAATAACGTACAAATGTGAAGAAAGCTTTGTGAAAATTCCTGGCGAGAAG
GACTCAGTGATCTGCCTTAAGGGCAGTCAATGGTCAGATATTGAAGAGTTCTGCAATCGTAGC
TGCGAGGTGCCAACAAGGCTAAATTCTGCATCCCTCAAACAGCCTTATATCACTCAGAATTAT
TTTCCAGTCGGTACTGTTGTGGAATATGAGTGCCGTCCAGGTTACAGAAGAGAACCTTCTCTA
TCACCAAAACTAACTTGCCTTCAGAATTTAAAATGGTCCACAGCAGTCGAATTTTGTAAAAAG
AAATCATGCCCTAATCCGGGAGAAATACGAAATGGTCAGATTGATGTACCAGGTGGCATATTA
TTTGGTGCAACCATCTCCTTCTCATGTAACACAGGGTACAAATTATTTGGCTCGACTTCTAGT
TTTTGTCTTATTTCAGGCAGCTCTGTCCAGTGGAGTGACCCGTTGCCAGAGTGCAGAGAAATT
TATTGTCCAGCACCACCACAAATTGACAATGGAATAATTCAAGGGGAACGTGACCATTATGGA
TATAGACAGTCTGTAACGTATGCATGTAATAAAGGATTCACCATGATTGGAGAGCACTCTATT
TATTGTACTGTGAATAATGATGAAGGAGAGTGGAGTGGCCCACCACCTGAATGC
TCGAGTCCTAACAAATGCACGCCTCCAAATGTGGAAAATGGAATATTGGTATCTGACAAC
AGAAGCTTATTTTCCTTAAATGAAGTTGTGGAGTTTAGGTGTCAGCCTGGCTTTGTCATG
AAAGGACCCCGCCGTGTGAAGTGCCAGGCCCTGAACAAATGGGAGCCGGAGCTACCAAGC
TGCTCCAGGGTATGTCAGCCACCTCCAGATGTCCTGCATGCTGAGCGTACCCAAAGGGAC
AAGGACAACTTTTCACCTGGGCAGGAAGTGTTCTACAGCTGTGAGCCCGGCTACGACCTC
AGAGGGGCTGCGTCTATGCGCTGCACACCCCAGGGAGACTGGAGCCCTGCAGCCCCCACA
TGTGAAGTGAAATCCTGTGATGACTTCATGGGCCAACTTCTTAATGGCCGTGTGCTATTT
CCAGTAAATCTCCAGCTTGGAGCAAAAGTGGATTTTGTTTGTGATGAAGGATTTCAATTA
AAAGGCAGCTCTGCTAGTTACTGTGTCTTGGCTGGAATGGAAAGCCTTTGGAATAGCAGT
GTTCCAGTGTGTGAACAAATCTTTTGTCCAAGTCCTCCAGTTATTCCTAATGGGAGACAC
ACAGGAAAACCTCTGGAAGTCTTTCCCTTTGGAAAAGCAGTAAATTACACATGCGACCCC
CACCCAGACAGAGGGACGAGCTTCGACCTCATTGGAGAGAGCACCATCCGCTGCACAAGT
GACCCTCAAGGGAATGGGGTTTGGAGCAGCCCTGCCCCTCGCTGTGGAATTCTGGGTCAC
TGTCAAGCCCCAGATCATTTTCTGTTTGCCAAGTTGAAAACCCAAACCAATGCATCTGAC
TTTCCCATTGGGACATCTTTAAAGTACGAATGCCGTCCTGAGTACTACGGGAGGCCATTC
TCTATCACATGTCTAGATAACCTGGTCTGGTCAAGTCCCAAAGATGTCTGTAAACGTAAA
TCATGTAAAACTCCTCCAGATCCAGTGAATGGCATGGTGCATGTGATCACAGACATCCAG
GTTGGATCCAGAATCAACTATTCTTGTACTACAGGGCACCGACTCATTGGTCACTCATCT
GCTGAATGTATCCTCTCGGGCAATGCTGCCCATTGGAGCACGAAGCCGCCAATTTGTCAA
CGAATTCCTTGTGGGCTACCCCCCACCATCGCCAATGGAGATTTCATTAGCACCAACAGA
GAGAATTTTCACTATGGATCAGTGGTGACCTACCGCTGCAATCCTGGAAGCGGAGGGAGA
AAGGTGTTTGAGCTTGTGGGTGAGCCCTCCATATACTGCACCAGCAATGACGATCAAGTG
GGCATCTGGAGCGGCCCGGCCCCTCAGTGCATTATACCTAACAAATGCACGCCTCCAAAT
GTGGAAAATGGAATATTGGTATCTGACAACAGAAGCTTATTTTCCTTAAATGAAGTTGTG
GAGTTTAGGTGTCAGCCTGGCTTTGTCATGAAAGGACCCCGCCGTGTGAAGTGCCAGGCC
CTGAACAAATGGGAGCCGGAGCTACCAAGCTGCTCCAGGGTATGTCAGCCACCTCCAGAT
GTCCTGCATGCTGAGCGTACCCAAAGGGACAAGGACAACTTTTCACCCGGGCAGGAAGTG
TTCTACAGCTGTGAGCCCGGCTATGACCTCAGAGGGGCTGCGTCTATGCGCTGCACACCC
CAGGGAGACTGGAGCCCTGCAGCCCCCACATGTGAAGTGAAATCCTGTGATGACTTCATG
GGCCAACTTCTTAATGGCCGTGTGCTATTTCCAGTAAATCTCCAGCTTGGAGCAAAAGTG

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GATTTTGTTTGTGATGAAGGATTTCAATTAAAAGGCAGCTCTGCTAGTTATTGTGTCTTG
GCTGGAATGGAAAGCCTTTGGAATAGCAGTGTTCCAGTGTGTGAACAAATCTTTTGTCCA
AGTCCTCCAGTTATTCCTAATGGGAGACACACAGGAAAACCTCTGGAAGTCTTTCCCTTT
GGAAAAGCAGTAAATTACACATGCGACCCCCACCCAGACAGAGGGACGAGCTTCGACCTC
ATTGGAGAGAGCACCATCCGCTGCACAAGTGACCCTCAAGGGAATGGGGTTTGGAGCAGC
CCTGCCCCTCGCTGTGGAATTCATCACCATCACCATCACTAAAGATCT

Fig. 9A

SEQ ID NO:15

M T V A R P S V P A A L P
L L G E L P R L L L L V L L C L P A V W G
D C G L P P D V P N A Q P A L E G R T S F
P E D T V I T Y K C E E S F V K I P G E K
D S V I C L K G S Q W S D I E E F C N R S
C E V P T R L N S A S L K Q P Y I T Q N Y
F P V G T V V E Y E C R P G Y R R E P S L
S P K L T C L Q N L K W S T A V E F C K K
K S C P N P G E I R N G Q I D V P G G I L
F G A T I S F S C N T G Y K L F G S T S S
F C L I S G S S V Q W S D P L P E C R E I
Y C P A P P Q I D N G I I Q G E R D H Y G
Y R Q S V T Y A C N K G F T M I G E H S I
Y C T V N N D E G E W S G P P P E C
S S P N K C T P P N V E N G I L V S D N
R S L F S L N E V V E F R C Q P G F V M
K G P R R V K C Q A L N K W E P E L P S
C S R V C Q P P P D V L H A E R T Q R D
K D N F S P G Q E V F Y S C E P G Y D L
R G A A S M R C T P Q G D W S P A A P T
C E V K S C D D F M G Q L L N G R V L F
P V N L Q L G A K V D F V C D E G F Q L
K G S S A S Y C V L A G M E S L W N S S
V P V C E Q I F C P S P P V I P N G R H
T G K P L E V F P F G K A V N Y T C D P
H P D R G T S F D L I G E S T I R C T S
D P Q G N G V W S S P A P R C G I L G H
C Q A P D H F L F A K L K T Q T N A S D
F P I G T S L K Y E C R P E Y Y G R P F
S I T C L D N L V W S S P K D V C K R K
S C K T P P D P V N G M V H V I T D I Q
V G S R I N Y S C T T G H R L I G H S S
A E C I L S G N A A H W S T K P P I C Q
R I P C G L P P T I A N G D F I S T N R
E N F H Y G S V V T Y R C N P G S G G R
K V F E L V G E P S I Y C T S N D D Q V
G I W S G P A P Q C I I P N K C T P P N

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V E N G I L V S D N R S L F S L N E V V
E F R C Q P G F V M K G P R R V K C Q A
L N K W E P E L P S C S R V C Q P P P D
V L H A E R T Q R D K D N F S P G Q E V
F Y S C E P G Y D L R G A A S M R C T P
Q G D W S P A A P T C E V K S C D D F M
G Q L L N G R V L F P V N L Q L G A K V
D F V C D E G F Q L K G S S A S Y C V L
A G M E S L W N S S V P V C E Q I F C P
S P P V I P N G R H T G K P L E V F P F
G K A V N Y T C D P H P D R G T S F D L
I G E S T I R C T S D P Q G N G V W S S
P A P R C G I L G H C Q A P D H F L F A
K L K T Q T N A S D F P I G T S L K Y E
C R P E Y Y G R P F S I T C L D N L V W
S S P K D V C K R K S C K T P P D P V N
G M V H V I T D I Q V
G S R I N Y S C T T G H R L I G H S S
A E C I L S G N A A H W S T K P P I C Q
R I P C G L P P T I A N G D F I S T N R
E N F H Y G S V V T Y R C N P G S G G R
K V F E L V G E P S I Y C T S N D D Q V
G I W S G P A P Q C I I P N K C T P P N
V E N G I L V S D N R S L F S L N E V V
E F R C Q P G F V M K G P R R V K C Q A
L N K W E P E L P S C S R V C Q P P P D
V L H A E R T Q R D K D N F S P G Q E V
F Y S C E P G Y D L R G A A S M R C T P
Q G D W S P A A P T C E V K S C D D F M
G Q L L N G R V L F P V N L Q L G A K V
D F V C D E G F Q L K G S S A S Y C V L
A G M E S L W N S S V P V C E Q I F C P
S P P V I P N G R H T G K P L E V F P F
G K A V N Y T C D P H P D R G T S F D L
I G E S T I R C T S D P Q G N G V W S S
P A P R C G I H H H H H H

Fig 9B
SEQ. ID NO:16

ATGACCGTCGCGCGGCCGAGCGTGCCCGCGGCGCTGCCC
CTCCTCGGGGAGCTGCCCCGGCTGCTGCTGCTGGTGCTGTTGTGCCTGCCGGCCGTGTGGGGT
GACTGTGGCCTTCCCCCAGATGTACCTAATGCCCAGCCAGCTTTGGAAGGCCGTACAAGTTTT
CCCGAGGATACTGTAATAACGTACAAATGTGAAGAAAGCTTTGTGAAAATTCCTGGCGAGAAG
GACTCAGTGATCTGCCTTAAGGGCAGTCAATGGTCAGATATTGAAGAGTTCTGCAATCGTAGC
TGCGAGGTGCCAACAAGGCTAAATTCTGCATCCCTCAAACAGCCTTATATCACTCAGAATTAT
TTTCCAGTCGGTACTGTTGTGGAATATGAGTGCCGTCCAGGTTACAGAAGAGAACCTTCTCTA

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TCACCAAAACTAACTTGCCTTCAGAATTTAAAATGGTCCACAGCAGTCGAATTTTGTAAAAAG
AAATCATGCCCTAATCCGGGAGAAATACGAAATGGTCAGATTGATGTACCAGGTGGCATATTA
TTTGGTGCAACCATCTCCTTCTCATGTAACACAGGGTACAAATTATTTGGCTCGACTTCTAGT
TTTTGTCTTATTTCAGGCAGCTCTGTCCAGTGGAGTGACCCGTTGCCAGAGTGCAGAGAAATT
TATTGTCCAGCACCACCACAAATTGACAATGGAATAATTCAAGGGGAACGTGACCATTATGGA
TATAGACAGTCTGTAACGTATGCATGTAATAAAGGATTCACCATGATTGGAGAGCACTCTATT
TATTGTACTGTGAATAATGATGAAGGAGAGTGGAGTGGCCCACCACCTGAATGC
TCGAGTCCTAACAAATGCACGCCTCCAAATGTGGAAAATGGAATATTGGTATCTGACAAC
AGAAGCTTATTTTCCTTAAATGAAGTTGTGGAGTTTAGGTGTCAGCCTGGCTTTGTCATG
AAAGGACCCCGCCGTGTGAAGTGCCAGGCCCTGAACAAATGGAGCCGGAGCTACCAAGC
TGCTCCAGGGTATGTCAGCCACCTCCAGATGTCCTGCATGCTGAGCGTACCCAAAGGGAC
AAGGACAACTTTTCACCTGGGCAGGAAGTGTTCTACAGCTGTGAGCCCGGCTACGACCTC
AGAGGGGCTGCGTCTATGCGCTGCACACCCCAGGGAGACTGGAGCCCTGCAGCCCCCACA
TGTGAAGTGAAATCCTGTGATGACTTCATGGGCCAACTTCTTAATGGCCGTGTGCTATTT
CCAGTAAATCTCCAGCTTGGAGCAAAAGTGGATTTTGTTTGTGATGAAGGATTTCAATTA
AAAGGCAGCTCTGCTAGTTACTGTGTCTTGGCTGGAATGGAAAGCCTTTGGAATAGCAGT
GTTCCAGTGTGTGAACAAATCTTTTGTCCAAGTCCTCCAGTTATTCCTAATGGGAGACAC
ACAGGAAAACCTCTGGAAGTCTTTCCCTTTGGAAAAGCAGTAAATTACACATGCGACCCC
CACCCAGACAGAGGGACGAGCTTCGACCTCATTGGAGAGAGCACCATCCGCTGCACAAGT
GACCCTCAAGGGAATGGGGTTTGGAGCAGCCCTGCCCCTCGCTGTGGAATTCTGGGTCAC
TGTCAAGCCCCAGATCATTTTCTGTTTGCCAAGTTGAAAACCCAAACCAATGCATCTGAC
TTTCCCATTGGGACATCTTTAAAGTACGAATGCCGTCCTGAGTACTACGGGAGGCCATTC
TCTATCACATGTCTAGATAACCTGGTCTGGTCAAGTCCCAAAGATGTCTGTAAACGTAAA
TCATGTAAAACTCCTCCAGATCCAGTGAATGGCATGGTGCATGTGATCACAGACATCCAG
GTTGGATCCAGAATCAACTATTCTTGTACTACAGGGCACCGACTCATTGGTCACTCATCT
GCTGAATGTATCCTCTCGGGCAATGCTGCCCATTGGAGCACGAAGCCGCCAATTTGTCAA
CGAATTCCTTGTGGGCTACCCCCCACCATCGCCAATGGAGATTTCATTAGCACCAACAGA
GAGAATTTTCACTATGGATCAGTGGTGACCTACCGCTGCAATCCTGGAAGCGGAGGGAGA
AAGGTGTTTGAGCTTGTGGGTGAGCCCTCCATATACTGCACCAGCAATGACGATCAAGTG
GGCATCTGGAGCGGCCCGGCCCCTCAGTGCATTATACCTAACAAATGCACGCCTCCAAAT
GTGGAAAATGGAATATTGGTATCTGACAACAGAAGCTTATTTTCCTTAAATGAAGTTGTG
GAGTTTAGGTGTCAGCCTGGCTTTGTCATGAAAGGACCCCGCCGTGTGAAGTGCCAGGCC
CTGAACAAATGGGAGCCGGAGCTACCAAGCTGCTCCAGGGTATGTCAGCCACCTCCAGAT
GTCCTGCATGCTGAGCGTACCCAAAGGGACAAGGACAACTTTTCACCCGGGCAGGAAGTG
TTCTACAGCTGTGAGCCCGGCTATGACCTCAGAGGGGCTGCGTCTATGCGCTGCACACCC
CAGGGAGACTGGAGCCCTGCAGCCCCCACATGTGAAGTGAAATCCTGTGATGACTTCATG
GGCCAACTTCTTAATGGCCGTGTGCTATTTCCAGTAAATCTCCAGCTTGGAGCAAAAGTG
GATTTTGTTTGTGATGAAGGATTTCAATTAAAAGGCAGCTCTGCTAGTTATTGTGTCTTG
GCTGGAATGGAAAGCCTTTGGAATAGCAGTGTTCCAGTGTGTGAACAAATCTTTTGTCCA
AGTCCTCCAGTTATTCCTAATGGGAGACACACAGGAAAACCTCTGGAAGTCTTTCCCTTT
GGAAAAGCAGTAAATTACACATGCGACCCCCACCCAGACAGAGGGACGAGCTTCGACCTC
ATTGGAGAGAGCACCATCCGCTGCACAAGTGACCCTCAAGGGAATGGGGTTTGGAGCAGC
CCTGCCCCTCGCTGTGGAATTCTGGGTCACTGTCAAGCCCCAGATCATTTTCTGTTTGCC
AAGTTGAAAACCCAAACCAATGCATCTGACTTTCCCATTGGGACATCTTTAAAGTACGAA
TGCCGTCCTGAGTACTACGGGAGGCCATTCTCTATCACATGTCTAGATAACCTGGTCTGG
TCAAGTCCCAAAGATGTCTGTAAACGTAAATCATGTAAAACTCCTCCAGATCCAGTGAAT
GGCATGGTGCATGTGATCACAGACATCCAGGTT
GGATCCAGAATCAACTATTCTTGTACTACAGGGCACCGACTCATTGGTCACTCATCT
GCTGAATGTATCCTCTCGGGCAATGCTGCCCATTGGAGCACGAAGCCGCCAATTTGTCAA
CGAATTCCTTGTGGGCTACCCCCCACCATCGCCAATGGAGATTTCATTAGCACCAACAGA
GAGAATTTTCACTATGGATCAGTGGTGACCTACCGCTGCAATCCTGGAAGCGGAGGGAGA

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AAGGTGTTTGAGCTTGTGGGTGAGCCCTCCATATACTGCACCAGCAATGACGATCAAGTG
GGCATCTGGAGCGGCCCGGCCCCTCAGTGCATTATACCTAACAAATGCACGCCTCCAAAT
GTGGAAAATGGAATATTGGTATCTGACAACAGAAGCTTATTTTCCTTAAATGAAGTTGTG
GAGTTTAGGTGTCAGCCTGGCTTTGTCATGAAAGGACCCCGCCGTGTGAAGTGCCAGGCC
CTGAACAAATGGGAGCCGGAGCTACCAAGCTGCTCCAGGGTATGTCAGCCACCTCCAGAT
GTCCTGCATGCTGAGCGTACCCAAAGGGACAAGGACAACTTTTCACCCGGGCAGGAAGTG
TTCTACAGCTGTGAGCCCGGCTATGACCTCAGAGGGGCTGCGTCTATGCGCTGCACACCC
CAGGGAGACTGGAGCCCTGCAGCCCCCACATGTGAAGTGAAATCCTGTGATGACTTCATG
GGCCAACTTCTTAATGGCCGTGTGCTATTTCCAGTAAATCTCCAGCTTGGAGCAAAAGTG
GATTTTGTTTGTGATGAAGGATTTCAATTAAAAGGCAGCTCTGCTAGTTATTGTGTCTTG
GCTGGAATGGAAAGCCTTTGGAATAGCAGTGTTCCAGTGTGTGAACAAATCTTTTGTCCA
AGTCCTCCAGTTATTCCTAATGGGAGACACACAGGAAAACCTCTGGAAGTCTTTCCCTTT
GGAAAAGCAGTAAATTACACATGCGACCCCCACCCAGACAGAGGGACGAGCTTCGACCTC
ATTGGAGAGAGCACCATCCGCTGCACAAGTGACCCTCAAGGGAATGGGGTTTGGAGCAGC
CCTGCCCCTCGCTGTGGAATTCATCACCATCACCATCACTAAAGATCT

Fig. 10A
SEQ. ID NO:17
ATA TAC GAA TTC TGG TTG AGT CCA AAT ATG GTC CC

Fig. 10B
SEQ. ID NO:18
ACA GTG AGA TCT TTA TCA TTT ACC CGG AGA CAG GGA G

Fig. 11A
SEQ. ID NO:19

M T V A R P S V P A A L P
L L G E L P R L L L L V L L C L P A V W G
D C G L P P D V P N A Q P A L E G R T S F
P E D T V I T Y K C E E S F V K I P G E K
D S V I C L K G S Q W S D I E E F C N R S
C E V P T R L N S A S L K Q P Y I T Q N Y
F P V G T V V E Y E C R P G Y R R E P S L
S P K L T C L Q N L K W S T A V E F C K K
K S C P N P G E I R N G Q I D V P G G I L
F G A T I S F S C N T G Y K L F G S T S S
F C L I S G S S V Q W S D P L P E C R E I
Y C P A P P Q I D N G I I Q G E R D H Y G
Y R Q S V T Y A C N K G F T M I G E H S I
Y C T V N N D E G E W S G P P P E C
S S P N K C T P P N V E N G I L V S D N

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R S L F S L N E V V E F R C Q P G F V M
K G P R R V K C Q A L N K W E P E L P S
C S R V C Q P P P D V L H A E R T Q R D
K D N F S P G Q E V F Y S C E P G Y D L
R G A A S M R C T P Q G D W S P A A P T
C E V K S C D D F M G Q L L N G R V L F
P V N L Q L G A K V D F V C D E G F Q L
K G S S A S Y C V L A G M E S L W N S S
V P V C E Q I F C P S P P V I P N G R H
T G K P L E V F P F G K A V N Y T C D P
H P D R G T S F D L I G E S T I R C T S
D P Q G N G V W S S P A P R C G I L
V E S K Y G P P C P S C P A P E F L
G G P S V F L F P P K P K D T L M I S R
T P E V T C V V V D V S Q E D P E V Q F
N W Y V D G V E V H N A K T K P R E E Q
F N S T Y R V V S V L T V L H Q D W L N
G K E Y K C K V S N K G L P S S I E K T
I S K A K G Q P R E P Q V Y T L P P S Q
E E M T K N Q V S L T C L V K G F Y P S
D I A V E W E S N G Q P E N N Y K T T P
P V L D S D G S F F L Y S R L T V D K S
R W Q E G N V F S C S V M H E A L H N H
Y T Q K S L S L S P G K

Fig. 11B
SEQ. ID NO:20

ATGACCGTCGCGCGGCCGAGCGTGCCCGCGGCGCTGCCC
CTCCTCGGGGAGCTGCCCCGGCTGCTGCTGCTGGTGCTGTTGTGCCTGCCGGCCGTGTGGGGT
GACTGTGGCCTTCCCCCAGATGTACCTAATGCCCAGCCAGCTTTGGAAGGCCGTACAAGTTTT
CCCGAGGATACTGTAATAACGTACAAATGTGAAGAAAGCTTTGTGAAAATTCCTGGCGAGAAG
GACTCAGTGATCTGCCTTAAGGGCAGTCAATGGTCAGATATTGAAGAGTTCTGCAATCGTAGC
TGCGAGGTGCCAACAAGGCTAAATTCTGCATCCCTCAAACAGCCTTATATCACTCAGAATTAT
TTTCCAGTCGGTACTGTTGTGGAATATGAGTGCCGTCCAGGTTACAGAAGAGAACCTTCTCTA
TCACCAAAACTAACTTGCCTTCAGAATTTAAAATGGTCCACAGCAGTCGAATTTTGTAAAAAG
AAATCATGCCCTAATCCGGGAGAAATACGAAATGGTCAGATTGATGTACCAGGTGGCATATTA
TTTGGTGCAACCATCTCCTTCTCATGTAACACAGGGTACAAATTATTTGGCTCGACTTCTAGT
TTTTGTCTTATTTCAGGCAGCTCTGTCCAGTGGAGTGACCCGTTGCCAGAGTGCAGAGAAATT
TATTGTCCAGCACCACCACAAATTGACAATGGAATAATTCAAGGGGAACGTGACCATTATGGA
TATAGACAGTCTGTAACGTATGCATGTAATAAAGGATTCACCATGATTGGAGAGCACTCTATT
TATTGTACTGTGAATAATGATGAAGGAGAGTGGAGTGGCCCACCACCTGAATGC
TCGAGTCCTAACAAATGCACGCCTCCAAATGTGGAAAATGGAATATTGGTATCTGACAAC
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AAGGACAACTTTTCACCTGGGCAGGAAGTGTTCTACAGCTGTGAGCCCGGCTACGACCTC

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AGAGGGGCTGCGTCTATGCGCTGCACACCCCAGGGAGACTGGAGCCCTGCAGCCCCCACA
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GAGGAGATGACCAAGAACCAGGTCAGCCTGACCTGCCTGGTCAAAGGCTTCTACCCCAGC
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CCCGTGCTGGACTCCGACGGCTCCTTCTTCCTCTACAGCAGGCTAACCGTGGACAAGAGC
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TACACACAGAAGAGCCTCTCCCTGTCTCCGGGTAAATGATAAAGATCT

Fig. 12A

SEQ. ID NO:21
ATA TAC GAA TTC TGG GTC ACT GTG AGG AGC CAC CAA CAT TTG AAG C

Fig. 12B

SEQ. ID NO:22
ACA GTG AGA TCT TTA GTG ATG GTG ATG GTG ATG CGA CAC TTT AAG ACA CTT TGG AAC

Fig. 13A

SEQ. ID NO:23

M	T	V	A	R	P	S	V	P	A	A	L	P							
L	L	G	E	L	P	R	L	L	L	L	V	L	L	C	L	P	A	V	W	G
D	C	G	L	P	P	D	V	P	N	A	Q	P	A	L	E	G	R	T	S	F
P	E	D	T	V	I	T	Y	K	C	E	E	S	F	V	K	I	P	G	E	K
D	S	V	I	C	L	K	G	S	Q	W	S	D	I	E	E	F	C	N	R	S
C	E	V	P	T	R	L	N	S	A	S	L	K	Q	P	Y	I	T	Q	N	Y
F	P	V	G	T	V	V	E	Y	E	C	R	P	G	Y	R	R	E	P	S	L
S	P	K	L	T	C	L	Q	N	L	K	W	S	T	A	V	E	F	C	K	K
K	S	C	P	N	P	G	E	I	R	N	G	Q	I	D	V	P	G	G	I	L
F	G	A	T	I	S	F	S	C	N	T	G	Y	K	L	F	G	S	T	S	S
F	C	L	I	S	G	S	S	V	Q	W	S	D	P	L	P	E	C	R	E	I

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Y C P A P P Q I D N G I I Q G E R D H Y G
Y R Q S V T Y A C N K G F T M I G E H S I
Y C T V N N D E G E W S G P P P E C
S S P N K C T P P N V E N G I L V S D N
R S L F S L N E V V E F R C Q P G F V M
K G P R R V K C Q A L N K W E P E L P S
C S R V C Q P P P D V L H A E R T Q R D
K D N F S P G Q E V F Y S C E P G Y D L
R G A A S M R C T P Q G D W S P A A P T
C E V K S C D D F M G Q L L N G R V L F
P V N L Q L G A K V D F V C D E G F Q L
K G S S A S Y C V L A G M E S L W N S S
V P V C E Q I F C P S P P V I P N G R H
T G K P L E V F P F G K A V N Y T C D P
H P D R G T S F D L I G E S T I R C T S
D P Q G N G V W S S P A P R C G I L G H
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E I G E R V D Y K C K K G Y F Y I P P L
A T H T I C D R N H T W L P V S D D A C
Y R E T C P Y I R D P L N G Q A V P A N
G T Y E F G Y Q M H F I C N E G Y Y L I
G E E I L Y C E L K G S V A I W S G K P
P I C E K V L C T P P P K I K N G K H T
F S E V E V F E Y L D A V T Y S C D P A
P G P D P F S L I G E S T I Y C G D N S
V W S R A A P E C K V V K C R F P V V E
N G K Q I S G F G K K F Y Y K A T V M F
E C D K G F Y L D G S D T I V C D S N S
T W D P P V P K C L K V S H H H H H H

Fig. 13B

SEQ. ID NO:24

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GACTCAGTGATCTGCCTTAAGGGCAGTCAATGGTCAGATATTGAAGAGTTCTGCAATCGTAGC
TGCGAGGTGCCAACAAGGCTAAATTCTGCATCCCTCAAACAGCCTTATATCACTCAGAATTAT
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ACTTGGGATCCCCCAGTTCCAAAGTGTCTTAAA//GTGTCG//CATCACCATCACCATCAC
TAAAGATCT

WESTERN BLOT OF HYBRID PROTEINS
DAF-IgG4, DAF-CR1BB, and DAF-CR1B

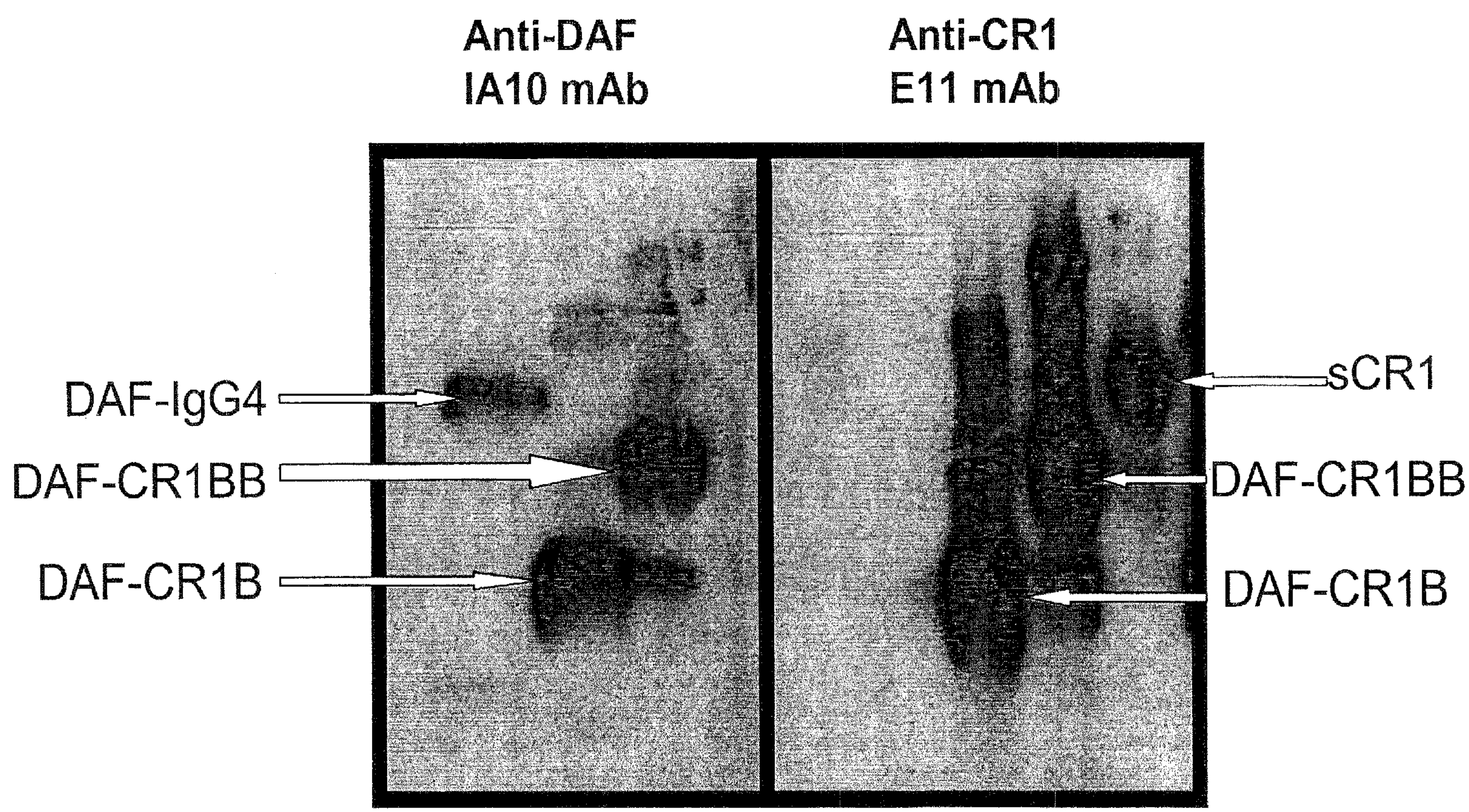

Fig. 14

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Western Blot of DAF-MCP

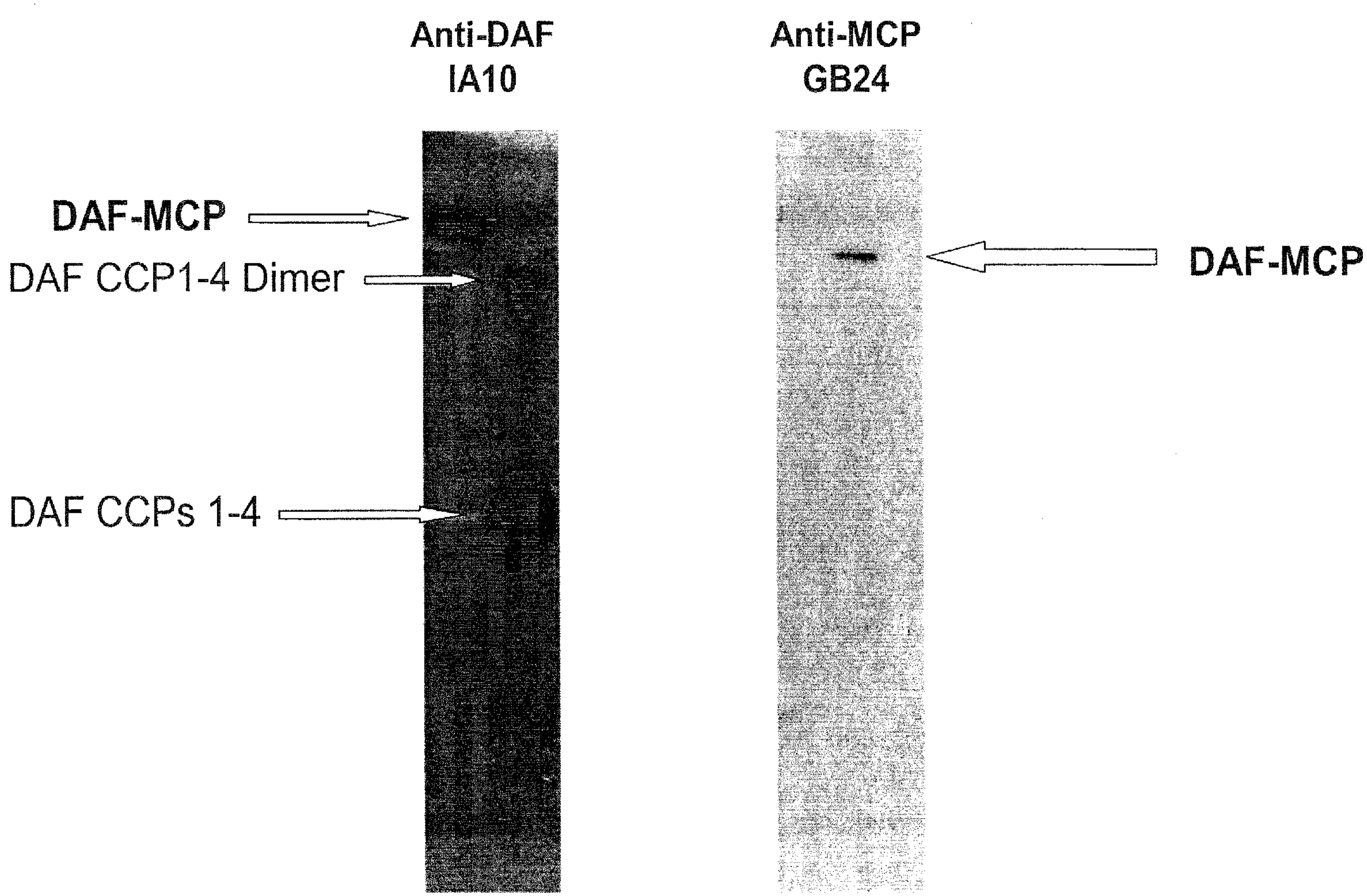

Fig. 15

Whole Serum Hemolytic Assay
DAF-CR1BB vs. Soluble CR1

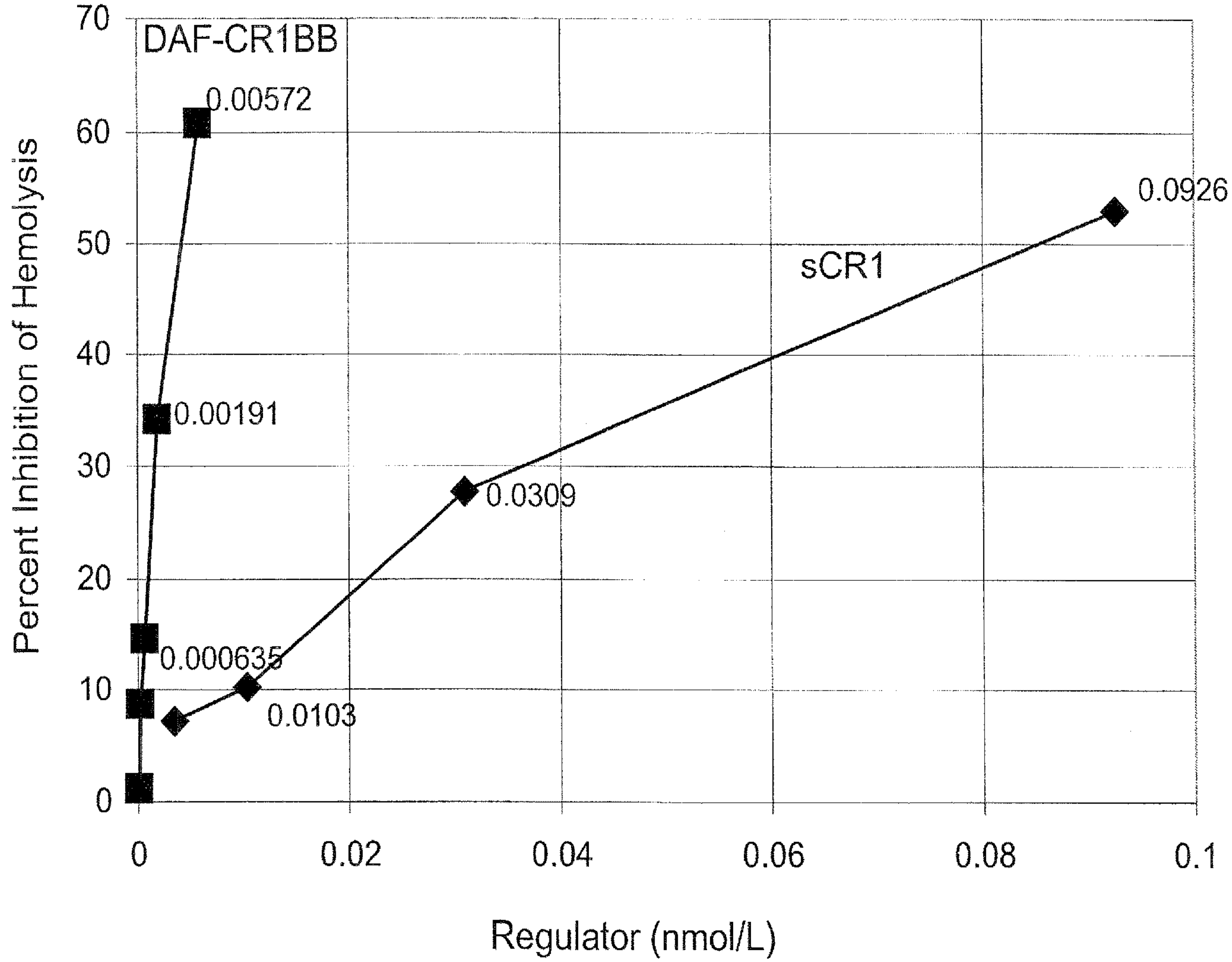

Fig. 16

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Whole Serum Hemolytic Assay
DAF-MCP Hybrid vs. DAF CCPs 1-4

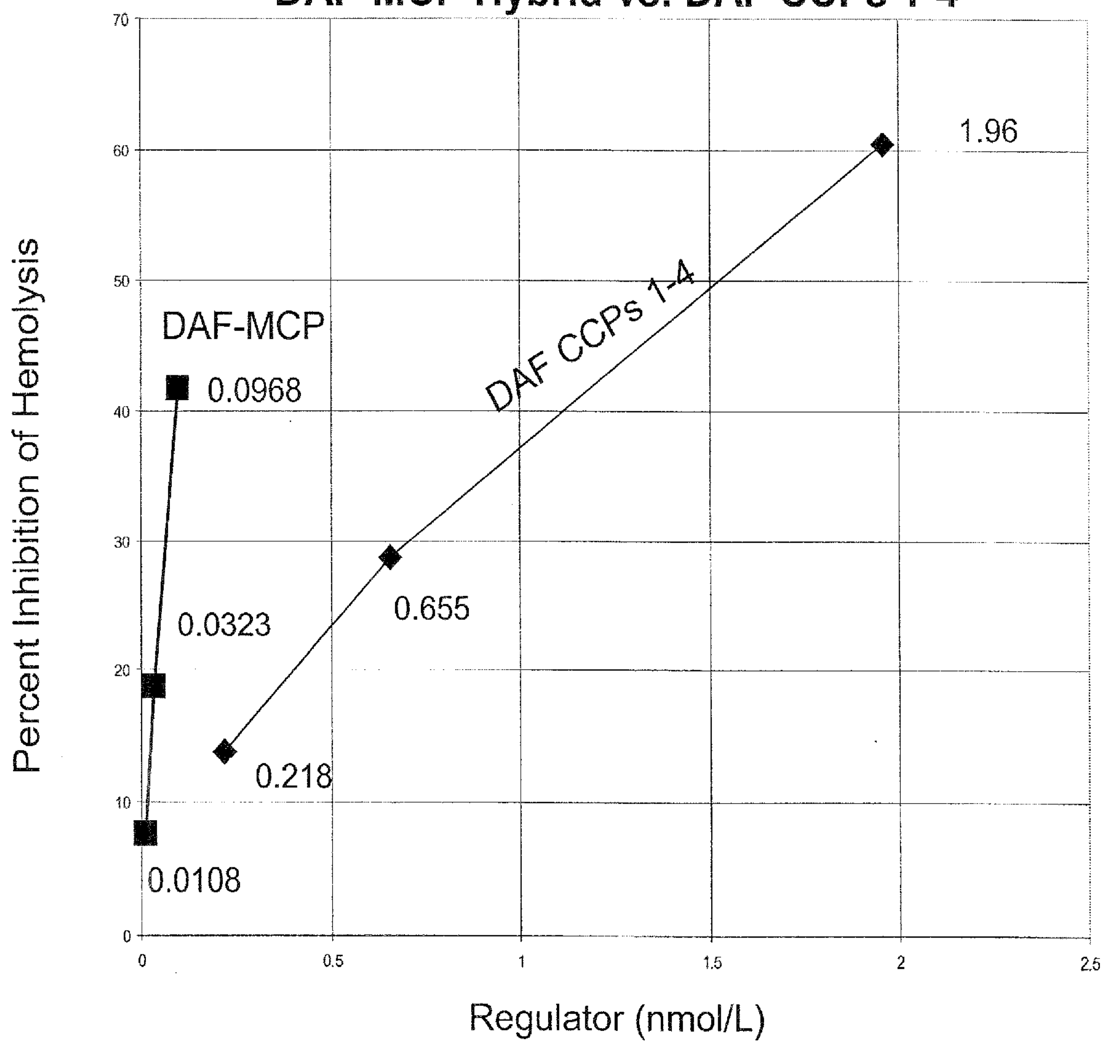

Fig. 17

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Classical Pathway C3 Convertase Decay

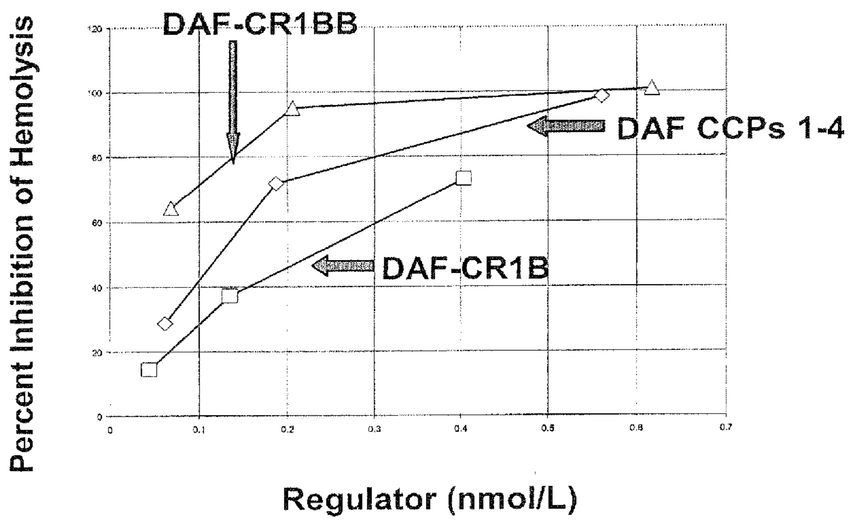

Fig. 18A

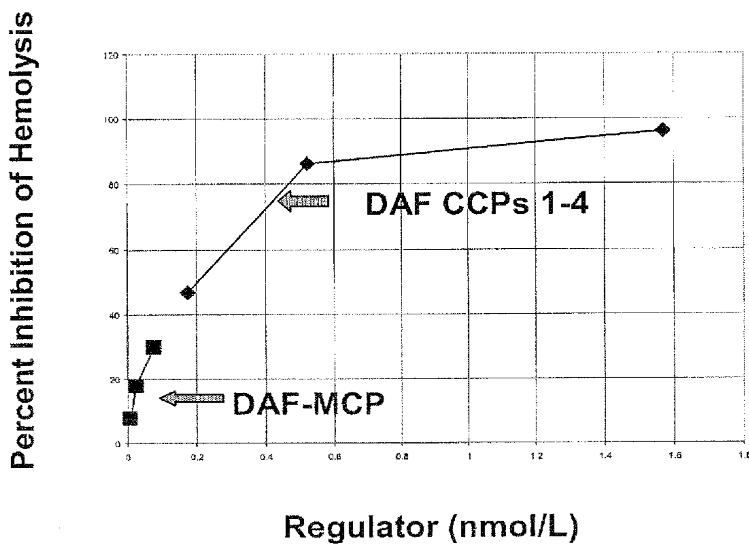

Fig. 18B

Classical Pathway C5 Convertase Decay
DAF-CR1B vs DAF CCPs 1-4

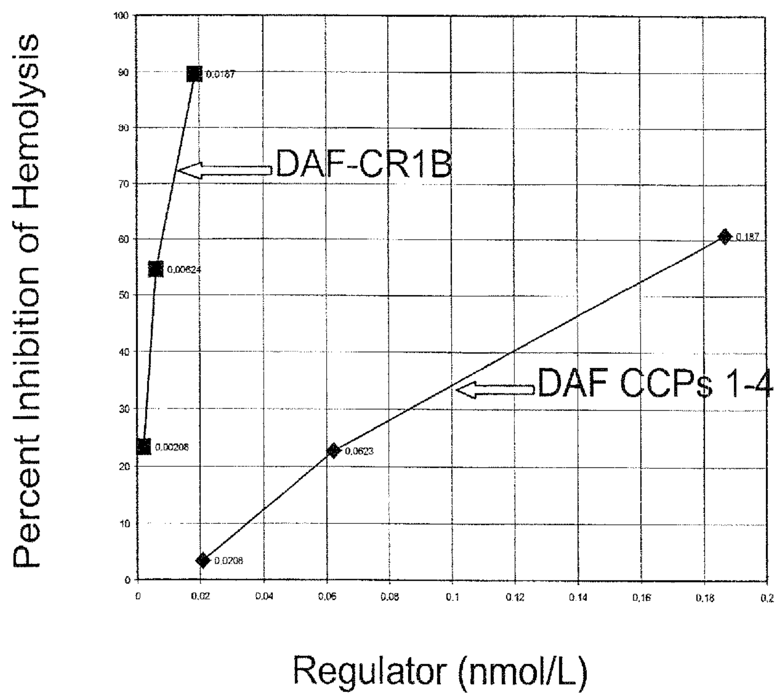

Fig. 19

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Classical Pathway C5 Convertase Decay
DAF-CR1BB vs sCR1 vs DAF-CR1B

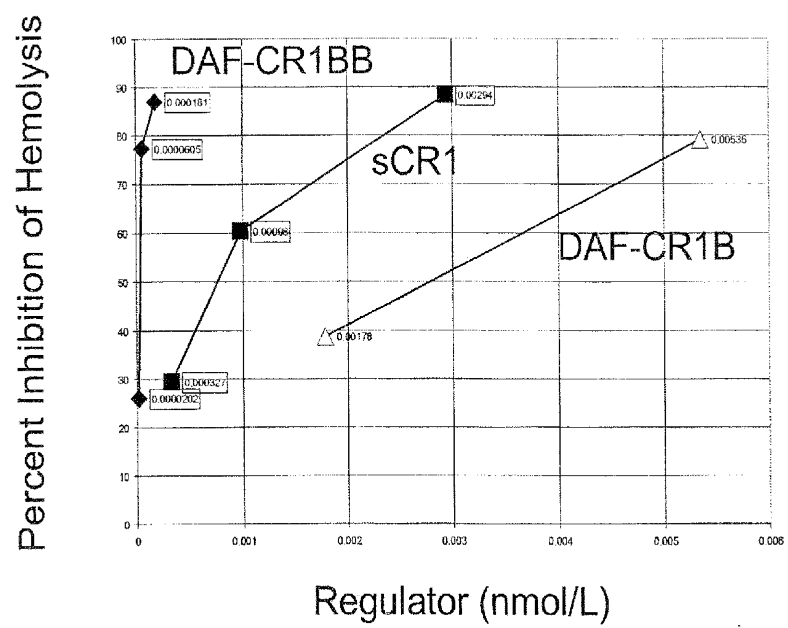

Fig. 20

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Cell-bound ($E^{sh}C4b3b$) Cofactor Assay

Cell Supernatant
(Anti-human C3 pAb)

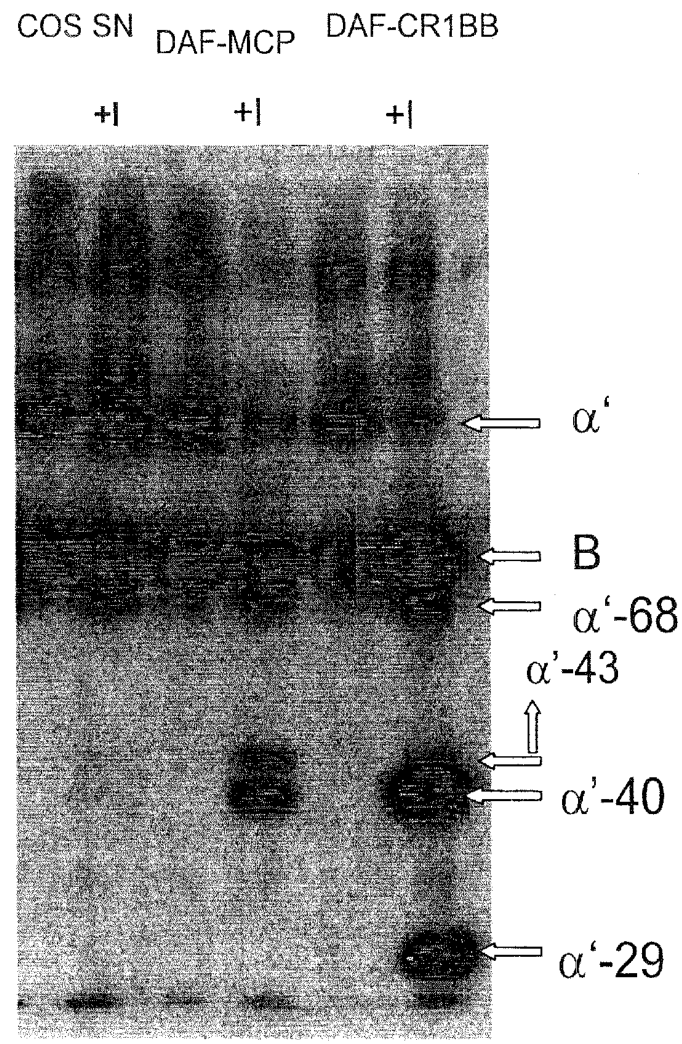

Fig. 21

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Cell-bound (EshC4b3b) Cofactor Assays
Cell Supernatant

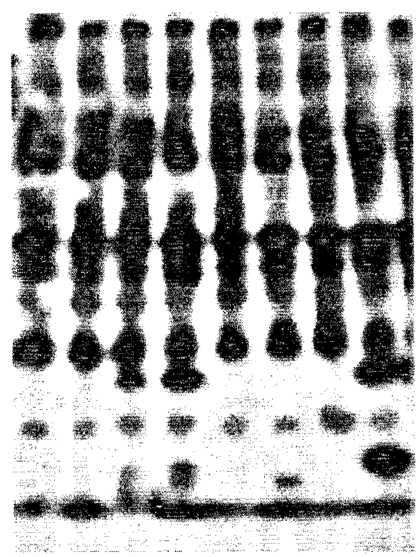

Fig. 22